RAW SEQUENCE LISTING
ERROR REPORT

BIOTECHNOLOGY
SYSTEMS
BRANCH

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10 517 214
Source: PCT
Date Processed by STIC: 12-22-04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
    http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
    U.S. Patent and Trademark Office, 220 20th Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lobby, Room 1B03, Arlington, VA 22202

Revised 05/17/04

PCT

RAW SEQUENCE LISTING   DATE: 12/22/2004
PATENT APPLICATION: US/10/517,214   TIME: 09:16:36

Input Set : **A:\3065 USOP seq.txt**
Output Set: **N:\CRF4\12222004\J517214.raw**

```
    2 <110> APPLICANT: MAEKAWA, Tsuyoshi
    3         HARA, Ryoma
    4         ODAKA, Hiroyuki
    5         MIZUFUNE, Hideya
    6         FUKATSU, Kohji
    8 <120> TITLE OF INVENTION: 1,2-Azole Derivatives with Hypoglycemic and Hypolipidemic
Activity
   10 <130> FILE REFERENCE: 3065 USOP
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/517,214
C--> 12 <141> CURRENT FILING DATE: 2004-11-23
   12 <150> PRIOR APPLICATION NUMBER: PCT/JP03/06389
   13 <151> PRIOR FILING DATE: 2003-05-22
   15 <150> PRIOR APPLICATION NUMBER: JP 2002-151405
   16 <151> PRIOR FILING DATE: 2002-05-24
   18 <150> PRIOR APPLICATION NUMBER: JP 2002-287161
   19 <151> PRIOR FILING DATE: 2002-09-30
   21 <150> PRIOR APPLICATION NUMBER: JP 2003-016748
   22 <151> PRIOR FILING DATE: 2003-01-24
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   26 <210> SEQ ID NO: 1
   27 <211> LENGTH: 34
   28 <212> TYPE: DNA
   29 <213> ORGANISM: Artificial Sequence
W--> 30 <220> FEATURE:
   31 <223> OTHER INFORMATION: Primer/probe
   33 <400> SEQUENCE: 1
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   36 <210> SEQ ID NO: 2
   37 <211> LENGTH: 34
   38 <212> TYPE: DNA
   39 <213> ORGANISM: Artificial Sequence
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   48 <212> TYPE: DNA
   49 <213> ORGANISM: Artificial Sequence
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   53 <400> SEQUENCE: 3
   54 ttagaattcg acatggacac caaacatttc ctg     33
   56 <210> SEQ ID NO: 4
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Does Not Comply
Corrected Diskette Needed

RAW SEQUENCE LISTING   DATE: 12/22/2004
PATENT APPLICATION: US/10/517,214   TIME: 09:16:36

                 Input Set : A:\3065 USOP seq.txt
                 Output Set: N:\CRF4\12222004\J517214.raw

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        58 <212> TYPE: DNA
        59 <213> ORGANISM: Artificial Sequence
W--> 60 <220> FEATURE:
        61 <223> OTHER INFORMATION: Primer/probe
        63 <400> SEQUENCE: 4
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        66 <210> SEQ ID NO: 5
        67 <211> LENGTH: 36
        68 <212> TYPE: DNA
        69 <213> ORGANISM: Artificial Sequence
W--> 70 <220> FEATURE:
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        73 <400> SEQUENCE: 5
        74 tcgacagggg accaggacaa aggtcacgtt cgggag 36
        76 <210> SEQ ID NO: 6
        77 <211> LENGTH: 36
        78 <212> TYPE: DNA
        79 <213> ORGANISM: Artificial Sequence
        81 <220> FEATURE:
        82 <223> OTHER INFORMATION:
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        87 <210> SEQ ID NO: 7
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        89 <212> TYPE: DNA
        90 <213> ORGANISM: Artificial Sequence
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        94 <400> SEQUENCE: 7
        95 cccagatctc cccagcgtct tgtcattg    28
        97 <210> SEQ ID NO: 8
        98 <211> LENGTH: 28
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       100 <213> ORGANISM: Artificial Sequence
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       104 <400> SEQUENCE: 8
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       108 <211> LENGTH: 33
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       114 <400> SEQUENCE: 9
       115 gtgggtaccg aaatgaccat ggttgacaca gag    33
       117 <210> SEQ ID NO: 10
       118 <211> LENGTH: 33
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[handwritten annotation:] pls explain source of genetic material. See item # 11 on error summary sheet.

[handwritten annotation near lines 79-84:] pls insert →

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        120 <213> ORGANISM: Artificial Sequence
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        129 <212> TYPE: DNA
        130 <213> ORGANISM: Artificial Sequence
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        134 <400> SEQUENCE: 11
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        137 <210> SEQ ID NO: 12
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W-->    146 ??
W-->    148 ??
W-->    150 ??
W-->    152 ??
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(handwritten annotation: circled lines 146–152 with "Pls delete")

VERIFICATION SUMMARY   DATE: 12/22/2004
PATENT APPLICATION: US/10/517,214   TIME: 09:16:37

Input Set : **A:\3065 USOP seq.txt**
Output Set: **N:\CRF4\12222004\J517214.raw**

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L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:30 M:283 W: Missing Blank Line separator, <220> field identifier
L:40 M:283 W: Missing Blank Line separator, <220> field identifier
L:50 M:283 W: Missing Blank Line separator, <220> field identifier
L:60 M:283 W: Missing Blank Line separator, <220> field identifier
L:70 M:283 W: Missing Blank Line separator, <220> field identifier
L:84 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:6,Line#:82
L:91 M:283 W: Missing Blank Line separator, <220> field identifier
L:101 M:283 W: Missing Blank Line separator, <220> field identifier
L:111 M:283 W: Missing Blank Line separator, <220> field identifier
L:121 M:283 W: Missing Blank Line separator, <220> field identifier
L:131 M:283 W: Missing Blank Line separator, <220> field identifier
L:141 M:283 W: Missing Blank Line separator, <220> field identifier
L:146 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:148 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:12
L:148 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:150 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:12
L:150 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:152 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:12
L:152 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
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STATISTICS SUMMARY   DATE: 12/22/2004
PATENT APPLICATION: US/10/517,214   TIME: 09:16:37

Input Set : **A:\3065 USOP seq.txt**
Output Set: **N:\CRF4\12222004\J517214.raw**

Application Serial Number: US/10/517,214
Alpha or Numeric or Xml: Numeric
Application Class:
Application File Date: 11-23-2004
Art Unit: PCT
Software Application:
Total Number of Sequences: 12
Total Nucleotides: 394
Total Amino Acids: 0
Number of Errors: 0
Number of Warnings: 19
Number of Corrections: 2

MESSAGE SUMMARY

258 W: 1 (Mandatory Feature missing)
270 C: 1 (Current Application Number differs)
271 C: 1 (Current Filing Date differs)
283 W: 11 (Missing Blank Line separator)
334 W: 4 ((2) Invalid Amino Acid in Coding Region)
336 W: 3 (Invalid Amino Acid Number in Coding Region)